

OIPE

RAW SEQUENCE LISTING DATE: 07/08/2002 PATENT APPLICATION: US/10/054,534A TIME: 10:34:30

Input Set : N:\Crf3\Refhold\J054534.raw
Output Set: N:\CRF3\07082002\J054534A.raw

P.D

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1 <110> APPLICANT: Abbott Laboratories
        Mukerji, Pradip
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         Huang, Yung-Sheng
         Das, Tapas
         Thurmond, Jennifer M.
         Pereira, Suzette L.
        Leonard, Amanda E.
 8 <120> TITLE OF INVENTION: DESATURASE GENES AND USES THEREOF
 9 <130> FILE REFERENCE: 6763.US.P1
10 <140> CURRENT APPLICATION NUMBER: US/10/054,534A
11 <141> CURRENT FILING DATE: 2002-05-21
12 <150> PRIOR APPLICATION NUMBER: US 09/769,863
13 <151> PRIOR FILING DATE: 2001-01-25
                                                              Does Not Comply
14 <160> NUMBER OF SEQ ID NOS: 55
                                                              Corrected Diskette Needed
15 <170> SOFTWARE: FastSEQ for Windows Version 4.0
17 <210> SEQ ID NO: 1
18 <211> LENGTH: 42
19 <212> TYPE: DNA
20 <213> ORGANISM: Artificial Sequence
21 <220> FEATURE:
22 <223> OTHER INFORMATION: Primer RO834
23 <220> FEATURE:
24 <221> NAME/KEY: misc_feature
25 <222> LOCATION: (3)...(3)
26 <223> OTHER INFORMATION: b = g or c or t/u at position 3
27 <220> FEATURE:
28 <221> NAME/KEY: misc_feature
29 <222> LOCATION: (6)...(6)
30 <223> OTHER INFORMATION: y = t/u or c at position 6
31 <220> FEATURE:
32 <221> NAME/KEY: misc_feature
33 <222> LOCATION: (9)...(9)
34 <223> OTHER INFORMATION: y = t/u or c at position 9
35 <220> FEATURE:
36 <221> NAME/KEY: misc_feature
37 <222> LOCATION: (12)...(12)
38 <223> OTHER INFORMATION: b = g or c or t/u at position 12
39 <220> FEATURE:
40 <221> NAME/KEY: misc_difference
41 <222> LOCATION: (18)...(18)
42 <223> OTHER INFORMATION: r = q or a at position 18
43 <220> FEATURE:
44 <221> NAME/KEY: misc_feature
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Input Set : N:\Crf3\Refhold\J054534.raw
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45 <222> LOCATION: (24)...(24) 46 <223> OTHER INFORMATION: b = g or c or t/u at position 24 47 <220> FEATURE: 48 <221> NAME/KEY: misc\_feature 49 <222> LOCATION: (30)...(30) 50 <223> OTHER INFORMATION: y = t/u or c at position 30 51 <220> FEATURE: 52 <221> NAME/KEY: misc\_feature 53 <222> LOCATION: (33)...(33) 54 <223> OTHER INFORMATION: y = t/u or c at position 33 55 <220> FEATURE: 56 <221> NAME/KEY: misc\_feature 57 <222> LOCATION: (36)...(36) 58 <223> OTHER INFORMATION: b = g or c or t/u at position 36 59 <220> FEATURE: 60 <221> NAME/KEY: misc\_feature 61 <222> LOCATION: (39)...(39) 62 <223> OTHER INFORMATION: h = a or c or t/u at position 39 63 <220> FEATURE: 64 <221> NAME/KEY: misc\_feature 65 <222> LOCATION: (42)...(42) 66 <223> OTHER INFORMATION: h = a or c or t/u at position 42 67 <400> SEQUENCE: 1 68 gtbtaygayg tbaccgartg ggtbaagcgy cayccbgghg gh 42 70 <210> SEQ ID NO: 2 71 <211> LENGTH: 45 72 <212> TYPE: DNA 73 <213> ORGANISM: Artificial Sequence 74 <220> FEATURE: 75 <223> OTHER INFORMATION: Forward Primer RO835 76 <220> FEATURE: 77 <221> NAME/KEY: misc\_feature 78 <222> LOCATION: (3)...(3) 79 <223> OTHER INFORMATION: h = a or c or t/u at position 3 80 <220> FEATURE: 81 <221> NAME/KEY: misc\_feature 82 <222> LOCATION: (6)...(6) 83 <223> OTHER INFORMATION: y = t/u or c at position 6 84 <220> FEATURE: 85 <221> NAME/KEY: misc\_feature 86 <222> LOCATION: (12)...(12) 87 <223> OTHER INFORMATION: y = t/u or c at position 12 88 <220> FEATURE: 89 <221> NAME/KEY: misc\_feature 90 <222> LOCATION: (27)...(27) 91 <223> OTHER INFORMATION: y = t/u or c at position 27 92 <220> FEATURE: 93 <221> NAME/KEY: misc\_feature 94 <222> LOCATION: (33)...(33)

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95 <223> OTHER INFORMATION: y = tu or c at position 33
96 <220> FEATURE:
                                                           'à is at location of 47
97 <221> NAME/KEY: misc_feature
98 <222> LOCATION: (39)...(39)
99 <223> OTHER INFORMATION: b = g or c or t/u at position 39
100 <220> FEATURE:
101 <221> NAME/KEY: misc_feature
102 <222> LOCATION: ((41)...(41)
103 <223> OTHER INFORMATION Y
                                 t/u or c at position 41
104 <220> FEATURE:
105 <221> NAME/KEY: misc_feature
106 <222> LOCATION: (45)...(45)
107 <223> OTHER INFORMATION: y = t/u or c at position 45
108 <400> SEQUENCE: 2
          gghgcytccg cyaactggtg gaagcaycag cayaacgtbc
                                                                             45
111 <210> SEQ ID NO: 3
112 <211> LENGTH: 45
113 <212> TYPE: DNA
114 <213> ORGANISM: Artificial Sequence
115 <220> FEATURE:
116 <223> OTHER INFORMATION: Reverse Primer RO836
117 <220> FEATURE:
118 <221> NAME/KEY: misc_feature
119 <222> LOCATION: (1)...(1)
120 <223> OTHER INFORMATION: r = g or a at position 1
121 <220> FEATURE:
122 <221> NAME/KEY: misc_feature
123 <222> LOCATION: (4)...(4)
124 <223> OTHER INFORMATION: r = g or a at position 4
125 <220> FEATURE:
126 <221> NAME/KEY: misc_feature
127 <222> LOCATION: (7)...(7)
128 <223> OTHER INFORMATION: v = a or g or c at position 7
129 <220> FEATURE:
130 <221> NAME/KEY: misc_feature
131 <222> LOCATION: (13)...(13)
132 <223> OTHER INFORMATION: r = q or a at position 13
133 <220> FEATURE:
134 <221> NAME/KEY: misc_feature
135 <222> LOCATION: (19)...(19)
136 <223> OTHER INFORMATION: r = g or a at position 19
137 <220> FEATURE:
138 <221> NAME/KEY: misc_feature
139 <222> LOCATION: (34)...(34)
140 < 223 > OTHER INFORMATION: r = q or a at position 34
141 <220> FEATURE:
142 <221> NAME/KEY: misc_feature
143 <222> LOCATION: (40)...(40)
144 <223> OTHER INFORMATION: r = g or a at position 40
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145 <220> FEATURE:
146 <221> NAME/KEY: misc_feature
147 <222> LOCATION: (43)...(43)
148 <223> OTHER INFORMATION: d = a or g or t/u at position 43
149 <400> SEQUENCE: 3
                                                                             45
          rtgrtgvacg ttrtgctgrt gcttccacca gttrgcggar gcdcc
150
152 <210> SEQ ID NO: 4
153 <211> LENGTH: 36
154 <212> TYPE: DNA
155 <213> ORGANISM: Artificial Sequence
156 <220> FEATURE:
157 <223> OTHER INFORMATION: Reverse Primer RO838
158 <220> FEATURE:
159 <221> NAME/KEY: misc_feature
160 <222> LOCATION: (6)...(6)
161 <223> OTHER INFORMATION: r = g or a at position 6
162 <220> FEATURE:
163 <221> NAME/KEY: misc_feature
164 <222> LOCATION: (12)...(12)
165 <223> OTHER INFORMATION: r = g or a at position 12
166 <220> FEATURE:
167 <221> NAME/KEY: misc_feature
168 <222> LOCATION: (15)...(15)
169 <223> OTHER INFORMATION: y = t/u or c at position 15
170 <220> FEATURE:
171 <221> NAME/KEY: misc_feature
172 <222> LOCATION: (18)...(18)
173 <223> OTHER INFORMATION: r = g or a at position 18
174 <220> FEATURE:
175 <221> NAME/KEY: misc_feature
176 <222> LOCATION: (21)...(21)
177 <223> OTHER INFORMATION: r = g or a at position 21
178 <220> FEATURE:
179 <221> NAME/KEY: misc_feature
180 <222> LOCATION: (24)...(24)
181 <223> OTHER INFORMATION: s = g or c at position 24
182 <220> FEATURE:
183 <221> NAME/KEY: misc_feature
184 <222> LOCATION: (27)...(27)
185 <223> OTHER INFORMATION: r = g or a at position 27
186 <220> FEATURE:
187 <221> NAME/KEY: misc_feature
188 <222> LOCATION: (30)...(30)
189 <223> OTHER INFORMATION: v = a or g or c at position 30
190 <400> SEQUENCE: 4
          ttgatrgtct arctygtrgt rgasaarggv tggtac
                                                                              36
191
193 <210> SEQ ID NO: 5
194 <211> LENGTH: 24
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195 <212> TYPE: DNA

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196 <213> ORGANISM: Artificial Sequence
     197 <220> FEATURE:
     198 <223> OTHER INFORMATION: Primer RO753
     199 <220> FEATURE:
     200 <221> NAME/KEY: misc_feature
     201 <222> LOCATION: (10)...(10)
     202 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown, or other at
              position 10
     203
     204 <220> FEATURE:
     205 <221> NAME/KEY: misc_feature
     206 <222> LOCATION: (13)...(13)
     207 <223> OTHER INFORMATION: r = g or a at position 13
     208 <220> FEATURE:
     209 <221> NAME/KEY: misc_feature
     210 <222> LOCATION: (16)...(16)
     211 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown, or other at
              position 16
     213 <220> FEATURE:
     214 <221> NAME/KEY: misc_feature
     215 <222> LOCATION: (18)...(19)
     216 <223> OTHER INFORMATION: r = g or a at positions 18-19
     217 <220> FEATURE:
     218 <221> NAME/KEY: misc_feature
     219 <222> LOCATION: (22)...(22)
     220 <223> OTHER INFORMATION: r = g or a at position 22
     221 <400> SEQUENCE: 5
                                                                                  24
               catcatcatn ggraanarrt grtg
W--> 222
     224 <210> SEQ ID NO: 6
     225 <211> LENGTH: 30
     226 <212> TYPE: DNA
     227 <213> ORGANISM: Artificial Sequence
     228 <220> FEATURE:
     229 <223> OTHER INFORMATION: Primer RO754
     230 <220> FEATURE:
     231 <221> NAME/KEY: misc_feature
     232 <222> LOCATION: (15)...(15)
     233 <223> OTHER INFORMATION: y = t/u or c at position 15
     234 <220> FEATURE:
     235 <221> NAME/KEY: misc_feature
     236 <222> LOCATION: (18)...(18)
     237 <223> OTHER INFORMATION: y = t/u or c at position 18
     238 <220> FEATURE:
     239 <221> NAME/KEY: misc_feature
     240 <222> LOCATION: (21)...(21)
     241 <223> OTHER INFORMATION: n = a or g or c or t/u, unknown, or other at
     242
               position 21
     243 <220> FEATURE:
     244 <221> NAME/KEY: misc_feature
     245 <222> LOCATION: (24)...(24)
```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/054,534A

DATE: 07/08/2002 TIME: 10:34:31

Input Set : N:\Crf3\Refhold\J054534.raw
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## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the  $\langle 220 \rangle$  to  $\langle 223 \rangle$  fields of each sequence which presents at least one n or Xaa.

Seq#:5; N Pos. 10,16
Seq#:6; N Pos. 21,27

VERIFICATION SUMMARY

DATE: 07/08/2002

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TIME: 10:34:31

Input Set : N:\Crf3\Refhold\J054534.raw
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L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0 L:257 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0